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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/721,212

DATE: 09/22/2003

TIME: 10:43:44

Input Set : A:\A-451K REV 091003-54.txt
 Output Set: N:\CRF4\09222003\I721212.raw

3 <110> APPLICANT: BOYLE, WILLIAM
 5 <120> TITLE OF INVENTION: OSTEOPROTEGERIN BINDING PROTEINS AND RECEPTORS
 7 <130> FILE REFERENCE: A-451K REV 09-10-03 54SEQ
 9 <140> CURRENT APPLICATION NUMBER: US 09/721,212
 10 <141> CURRENT FILING DATE: 2000-11-21
 12 <150> PRIOR APPLICATION NUMBER: US 09/052,521
 13 <151> PRIOR FILING DATE: 1998-03-30
 15 <150> PRIOR APPLICATION NUMBER: US 08/880,855
 16 <151> PRIOR FILING DATE: 1997-06-23
 18 <150> PRIOR APPLICATION NUMBER: US 08/842,842
 19 <151> PRIOR FILING DATE: 1997-04-16
 21 <160> NUMBER OF SEQ ID NOS: 54
 23 <170> SOFTWARE: PatentIn version 3.1
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 2295
 27 <212> TYPE: DNA
 28 <213> ORGANISM: Mus musculus
 30 <220> FEATURE:
 31 <221> NAME/KEY: CDS
 32 <222> LOCATION: (158)..(1105)
 33 <223> OTHER INFORMATION:
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 41 tcgcggagca gggcgccccga actccggcgcc atg cgc cgg gcc agc cga 175
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 43 1 5
 45 gac tac ggc aag tac ctg cgc agc tcg gag gag atg ggc agc ggc ccc 223
 46 Asp Tyr Gly Lys Tyr Leu Arg Ser Ser Glu Glu Met Gly Ser Gly Pro
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 49 ggc gtc cca cac gag ggt ccg ctg cac ccc gcg cct tct gca ccg gct 271
 50 Gly Val Pro His Glu Gly Pro Leu His Pro Ala Pro Ser Ala Pro Ala
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 62 Phe Arg Ala Gln Met Asp Pro Asn Arg Ile Ser Glu Asp Ser Thr His
 63 75 80 85
 65 tgc ttt tat aga atc ctg aga ctc cat gaa aac gca ggt ttg cag gac 463

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73	aaa	caa	gcc	ttt	cag	ggg	gcc	gtg	cag	aag	gaa	ctg	caa	cac	att	gtg	559	
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 143 aatatttaaa aatgtctcac tggtgacata tttatgttt taaatgtaca gatgtattt 1735
 145 actgggcac tttgttaattc ccctgaaggt actcgtagct aaggggcag aatactgttt 1795
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 149 gtcaaaaacta tgcaagcaaa ataaataat aaaaataaaa tgaataacctt gaataataag 1915
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 157 atagaaaatg tactatagtt ttcagccaa aaaacaatcc actttttat ttagtggaaag 2155
 159 ttat ttttatactgtaca ataaaagcat tgtctctgaa tgttaatttt ttggtaaaaa 2215
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 182 35 40 45
 185 Met Phe Leu Ala Leu Leu Gly Leu Gly Leu Gly Gln Val Val Cys Ser
 186 50 55 60
 189 Ile Ala Leu Phe Leu Tyr Phe Arg Ala Gln Met Asp Pro Asn Arg Ile
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 193 Ser Glu Asp Ser Thr His Cys Phe Tyr Arg Ile Leu Arg Leu His Glu
 194 85 90 95
 197 Asn Ala Gly Leu Gln Asp Ser Thr Leu Glu Ser Glu Asp Thr Leu Pro
 198 100 105 110
 201 Asp Ser Cys Arg Arg Met Lys Gln Ala Phe Gln Gly Ala Val Gln Lys
 202 115 120 125
 205 Glu Leu Gln His Ile Val Gly Pro Gln Arg Phe Ser Gly Ala Pro Ala
 206 130 135 140
 209 Met Met Glu Gly Ser Trp Leu Asp Val Ala Gln Arg Gly Lys Pro Glu
 210 145 150 155 160
 213 Ala Gln Pro Phe Ala His Leu Thr Ile Asn Ala Ala Ser Ile Pro Ser
 214 165 170 175
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 218 180 185 190
 221 Ala Lys Ile Ser Asn Met Thr Leu Ser Asn Gly Lys Leu Arg Val Asn
 222 195 200 205
 225 Gln Asp Gly Phe Tyr Tyr Leu Tyr Ala Asn Ile Cys Phe Arg His His
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 229 Glu Thr Ser Gly Ser Val Pro Thr Asp Tyr Leu Gln Leu Met Val Tyr

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